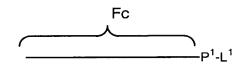


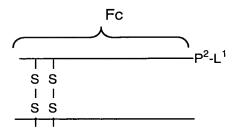
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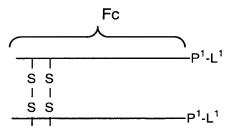
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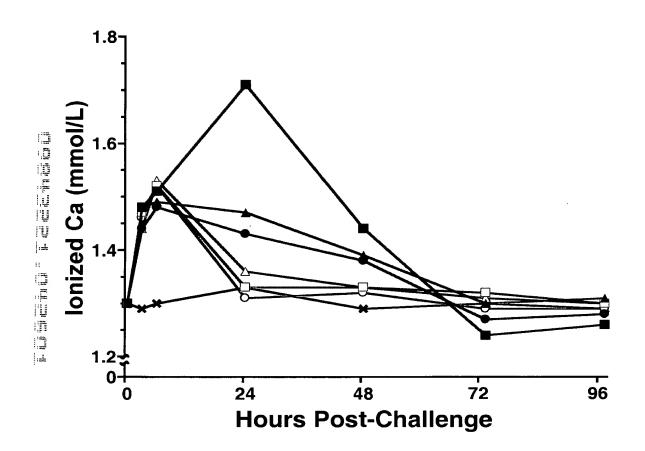
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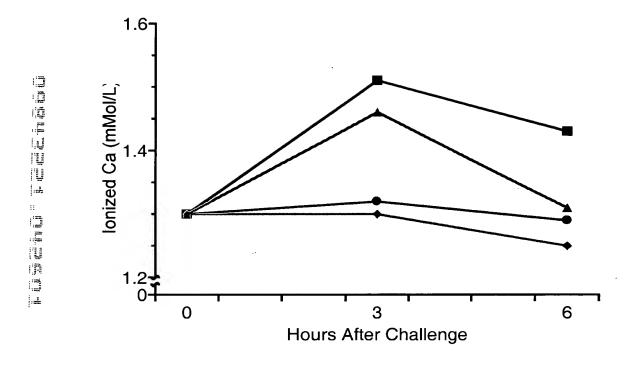
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P. C.

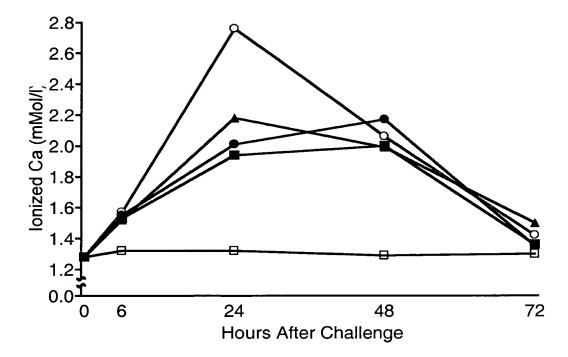
## FIG. 3

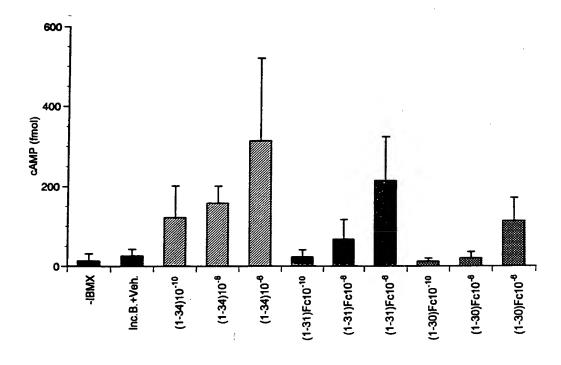
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	_	TACC																			. 60				
a		M D	K	т	н	т	С	P	P	С	P	A	P	E	L	L	G	G	P	s	-				
	61	GTCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTC															100								
			CAGAAGGAGAAGGGGGTTTTGGGTTCCTGTGGGAGTACTAGAGGGCCTGGGGACTCCAG														120								
a		V F	L	F	P	P	K	P	K	D	Т	L	М	I	s	R	Т	P	E	V	-				
	121	ACATGCGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAAGTTCAACTGGTACGTG														180									
		TGTACGCACCACCTGCACTCGGTGCTTCTGGGACTCCAGTTCAAGTTGACCATGCAC														100									
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	181	GACGO																			240				
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2	361	TTTCC	GCA  CGT	GCC -+- CGG	CCG.  GGC'	AGA.  TCT	ACC. + TGG	ACA  TGT	GGT(  CCA	GTA(	CAC( + GTG(	CCT(  GGA(	GCC(  CGG(	CCC -+- GGG'	ATC(  TAG(	CCG(  GGC(	GGA'	rgac	GAC	GACC + CTGG	420				
a	361	TTTCC	GGCA CGGT Q	GCC -+- CGG	CCG  GGC' R	AGA.  TCT E	ACC. + TGG	ACA  TGT	GGT(  CCA( V	GTAG  CATG	CACO + GTGO T	CCTC  GGAC L	GCC GGG P	CCC -+- GGG'	ATCO  FAGO S	CCG(  GGC( R	GGA' + CCTI	rgac  ACTC	GCTC GAC	GACC + CTGG	420				
a		TTTCC K G	GGCA CGGT Q	GCC -+- CGG P GGT	CCG GGC' R CAG	AGA TCT E CCT	ACC + TGG P GAC	ACA TGT Q CTG	GGT CCA V	GTAG CATG Y GGTG	CACO H GTGO T	CCT( GGA( L AGG(	GCCC CGGC P	CCC. -+- GGG' P CTA'	ATCO	CCGC GGCC R CAGC	D CGA	rgac ACTC E	GCTC CGAC L	GACC CTGG T	-				
		TTTCC	GGCA CGGT Q ACCA	GCC( -+- CGG( P GGT( -+- CCA(	CCG. GGC' R CAGG	AGA. TCT E CCT  GGA	ACC. TGG P GAC	ACA TGT Q CTG CTG	GGT CCA V CCT GGA	GTAG CATG Y GGTG	CACO H GTGO T CAA H GTT'	CCTC GGAC L AGGC	GCCC CGGC P CTTC	CCC. GGG' P CTA' -+	ATCO FAGO S FCCO AGGO	CCGC GGCC R CAGC	GGAGGGCTG	TGACTC E CATC	GCTC CGAC L CGCC	GACC TTGG	-				
a		TTTCC K G	GGCA CGGT Q ACCA CGGT	GCC -+- CGG P GGT -+- CCA	CCG. GGC'  R CAGG	AGA. TCT E CCT  GGA	ACC. TGG  P GACC+ CTGC	ACA TGT Q CTG  GAC	GGT CCA V CCT  GGA	GTAG CATG Y GGTG CCAG	CACC TCAA CAA + GTT	CCTC GGAC L AGGC TCCC	GCCC CGGC P CTTC GAAC	CCC. P CTA' -+ GAT.	ATCO FAGO S TCCO AGGO	CCGC GGCC R CAGC GTCC	D CGAG GCTG	TGACTC E CATC	EGCC EGCC L EGCC EGCC	GACC T CGTG CGTG GCAC	-				
	421	TTTCC  K G  AAGAA  TTCTT  K N	GGCA Q ACCA CGGT Q Q GGGA	GCC -+- CGG P GGT -+- CCA V GAG -+-	CCG. GGC'  R CAGG GTCG S CAA'	AGA. TCT E CCT GGA	ACC. TGG  P GAC CTG  T GCA+	ACA TGT Q CTG GAC	GGT CCA V CCT GGA L	GTAG CATG Y GGTG CCAG	CACC GTGC T CAAC + GTT' K	CCTO GGAO L AGGO TCCO	GCCC CGGC P CTTC GAAC F	CCC. GGG'  P  CTA' GAT.  GACC	ATCO PAGGO P	CCGC GGCC R CAGC GTCC	GGAC  CGAC  GCTC  D  CGAC  CGAC  CCTC	FGACTO E CATO	ECTO CGAO L CGCO GCGO	GACC T CGTG GCAC V GGAC	- 480 -				
	421	TTTCC  K G  AAGAA  TTCTT  K N  GAGTC	GGCA Q ACCA CGGT Q Q GGGA	GCC -+- CGG P GGT -+- CCA V GAG -+-	CCG. GGC'  R CAGG GTCG S CAA'	AGA. TCT E CCT GGA	ACC+ TGG'  GAC'+ CTG'  GCA'+ CGT'	ACA TGT Q CTG GAC	GGT CCA V CCT GGA L	GTAG CATG Y GGTG CCAG	CACC GTGC T CAAC + GTT' K	CCTO GGAO L AGGO TCCO	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC. GGG'  P  CTA' GAT.  GACC	ATCO PAGGO P	CCGC GGCC R CAGC GTCC	GGAC  CGAC	FGACTO E CATO	ECTO CGAO L CGCO A ECTO	GACC T CGTG GCAC V GGAC	- 480 -				
a	421 481	TTTCC  K G  AAGAA  TTCTT  K N  GAGTC  CTCAC	GGCA Q ACCA TGGT Q GGGA CCCT E	GCCCCACCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCG, GGC'  R CAGG S CAA' S CAA' N CTTC	AGA. TCT  E CCTCGGA  L TGGGC G CCTTC	ACC+ TGG P GAC+ CTG T GCA+ CGT.	ACAC TGT Q CTGG GACC C GCCG P	GGTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCT	GTA( CATC  Y  GGTY  CCA( V  GAA( CTTY  N  CCAA(	CACC + GTGG T CAAA K CAAG K CAAG N	CCTCGGAC	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC+ GGG  P CTA -+ GAT.  Y GAC+ CTGG	ATCC FIRGO S FICCO P CACO T	CCGC R CAGC S S S CCGGA	D CCGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	TGACTC  E CATCC  I CGGCAC  V	GCTCCGAC  L CCGCCCGAC  A  CCCGAC  L  CCGCCCCGAC  L  CCGCCCGAC  L  GCCCGAC	GACC T CGTG GCAC V GGAC CCTG D	- 480 - 540 -				
a	421 481	TTTCC  K G  AAGAA  TTCTT  K N  GAGTC  CTCAC	GGCA Q ACCA PGGT Q GGGA CCT E	GCCC -+- CGGG P GGTC-  CCA V GAGG-  CTCC S	CCG, GGC'  R CAGGTC  S CAA'  S CTTC	AGA. TCT  E CCT(GGA)  L TGGG G CTT(GGC) G	ACC+ TGG P GAC+ CTG T GCA+ CGT Q CCT	ACAC	GGTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCTCCCT	GTA( CATC  Y  GGTY  CCA  V  GAA( CTTY  N  CAA( CAA( CAA( CAA( CAA( CAA( CAA(	CACC + GTGG T CAAA K CCAAG K CCAAG N GCTG	CCTCGGAC	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC. SGG'  P CTA'  Y SACC CTGG  T	ATCC FAGG P CACC T	CCGCCGG	D CCGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	TGACTC E CATC I CCGTC V STGC	GCTCCGAC  L CCGCCCGAC  A L L CCGCCCGAC	GACC T CGTG GCAC V GGAC CCTG D GCAG	- 480 - 540 -				
a	421 481	TTTCO K G AAGAA TTCTT K N GAGTO CTCAC	GGCA Q ACCA CGGT Q ACCA CGGT Q CGGGCA CCCT E	GCCC -+- CGGG P GGTC -+- CCCA V GAGGG S CTCC GAGG	CCG. GGC'  R CAGG GTCG S CAA GTTC N CTTCGAA	AGA. TCT' E CCTGGGA L TGGGG G CCTTGGGAA	ACC+ TGG P GACC+ CTGG T GCAGC+ CGTG Q CCTGGGGA	ACAC	GGTC	GTA( Y  GGT( CCA( V  GAA( I I I I I I I I I I I I I I I I I I	CACO + GTG T CAAL GTT K CAAC GTT N GCTC CGAC	CCTCGGACCGGACCGGGGGGGGGGGGGGGGGGGGGGGGG	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC.  P CTA'  Y  GAC  T  GGAC  CCTC	ATCO	CCGGG	D CGA( D	TGAC E ECATC	EGCTC  L  CGGCC  A  GGCTC  L  CGGCCC  CGGCC  CGGCCC  CGGCC  CGCC  CGC	GACC T CGTG T CGTG V GGAC CCTG D GCAG	- 480 - 540 -				
a	421 481 -541-	TTTCO K G AAGAA TTCTT K N GAGTO CTCAO E W TCCGA AGGCT S D	GGCA  Q ACCA ACCA Q ACCA CGGT  Q ACCA CGGCC  E ACGG ACGCC G ACGCC	GCCCACCCCACCCCACCCCACCCCACCCCACCCCACCC	CCG. GGC'  R CAGG S CAA' N CTTC GAA F	AGA TCT E CCTGGA L TGGG G CTTGGG F GAACC	ACC+ TGG P GACC+ CTG T GCA+ CGT Q CCT+ GGA L CTC	ACA' Q Q CTGG GAC' C GCGG P CTA' Y CGTC	GGTO	GTA( Y  GGTCCA( V  GAA( CTTCCA  N  CAA( GTTCCA  K  GCA( GCA( GCA( GCA( GCA( GCA( GCA( GCA	CACC + GTGC T CAAAC K CAAC N GCTC N GCTC CGAC	CCTC GGAC CTAC GCTAC Y CACC GTGC T CCC T C CCC T C C C C C C C C C C C C C C C C C C C C	F CAAC	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	ATCO S FCCC P CACC T CAACC T CAACC K K CAACC	CCGGG	D CGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	TGAC	GCTCCGAC  L CGCCGAC  A GCTCCGAC  L GCCAC  L GCCAC  Q	GACC T CGTG GCAC V GGAC CCTG D GCAG CGTC Q GAAG	- 480 - 540 - - -6.0.0_				
a	421 481 -541-	TTTCO  K G  AAGAA  TTCTT  K N  GAGTO  CTCAC  E W  TCCGA  AGGCT  S D	GGCA  Q ACCA ACCA Q GGGGA  CCCT E ACGG GGCC G	GCCC -+- CGGG P GGTC CCCA V GAGGG S CTCC S CTCC S CTCC -+- GAGG S	CCG. GGC' R CAGG S S CAA' GTTC GTTC F CCTC CCTC CCTC CCTC CCTC CC	AGA  TCT  E  CCTC  GGA  L  TGGC  G  CCTTC  GAAC  F  ATGG	ACC+ TGG P GAC'+ CTG CTG CCT' Q CCT' GGA' L CTC'	ACA' Q CTGC GACC C GCCGG P CTAC GATC	GGTC	GTAG  Y  GGTG  V  GAAG  CTTG  N  CAAG  K  GCA	CACC TTCAAL KCAAC STTC K CAAC CCAC L	CCTC  L AGGG GTAC  GTAC  GTAC  T CACC  T GGGTG  T	GCCCC P CTTCC GAAC F CAAC K CCGTCC V CCTCC V	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	ATCO FACO PCACO TCAACO TCAACO K	CCGGC R CCAGGCC S S S CCGGGA	D CCGA( D PCCCC P CAGG R R CCTAC	TGAC	EGCTC  L  CCGCC  A  GCCCGAC  L  CCCGAC  Q  GCCAC	GACC T CGTG CGTG CGTG V GGAC CCTG D GCAG CGTC Q GAAG	- 480 - 540 - - -6.0.0_				
a	421 481 -541-	TTTCO  K G  AAGAA  TTCTT  K N  GAGTO  CTCAC  E W  TCCGA  AGGCT  S D  GGGAA  CCCTT	GGCA  Q ACCA ACCA Q GGGGA  CCCT E ACGG GGCC G	GCCCCACCCCACCCCCACCCCCCCCCCCCCCCCCCCCC	CCG. R CAGGC S CAAA CTTC GAA F CTCC GAGG	AGA TCT E CCTTCGGA CTTCGGAA F ATGC	ACC+ TGG P GAC+ CTG T GCA+ CGT Q CCT+ GGA L CTC. GAG	ACA	GGTCCCTAC	GTAGE CATO	CAC( T GTG  T CAAL + GTT'  K CAAC T GTTC  L TGAC  L TGAC  TGAC CCC  TGAC  TG	CCT(CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC. P CTA -+- GAT: Y GAC: -+- CCTG D GCA: CCTGT(	ATCO FINE S FICCO AGGO FINE STEEL T CAAC K K CAAC	CCGGC R CAGCC S S GCCC P GAGCC S GCCCGGA	D CGACO P CAGO R R CTACO R CTA	TGAC E ECATC I CGTC V GTGC W W CACC	GCTC  CGAC  L  CGGCGC  A  GCTC  CGAC  Q  GCAC	GACC T CGTG CGTG CGTG V GGAC CCTG D GCAG CGTC Q GAAG	- 480 - 540 - - -6.0.0_				
a a	421 481 -541- 601	TTTCO K G AAGAA TTCTT K N GAGTO CTCAO E W TCCGA AGGCT G N AGCCTT	GGCA  Q ACCA ACCA Q GGGA  CCCT  E ACGG GCA CCCT  CCCT	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCG. GGC' R CAGG S S CAA' GTTC GTTC F CTTC GGAG F CTTC GGAG S GGGG S GGGG S GGGG S	AGA  E CCTTC GGA  L TGGG G CTTC GAAC  F ATGG CTTAC C TTCC	ACC+ TGG P GAC'+ CTG T GCA' CGT' CGT' CGAG S GGG'	ACA- TGT- Q CTG- GAC C GGCC- CGGG P CTA- Y CGTY V TAA	GGTCCTAC	GTA( Y GGT( CCA( V GAA( CTT( K GCA  K GCA  H	CAC( T GTG  T CAAL + GTT'  K CAAC T GTTC  L TGAC  L TGAC  TGAC CCC  TGAC  TG	CCT(CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC. P CTA -+- SAT: Y SAC: -+- CCTG D GCA: CCTGT(	ATCO FINE S FICCO AGGO FINE STEEL T CAAC K K CAAC	CCGGC R CAGCC S S GCCC P GAGCC S GCCCGGA	D CGACO P CAGO R R CTACO R CTA	TGAC E ECATC I CGTC V GTGC W W CACC	GCTC  CGAC  L  CGGCGC  A  GCTC  CGAC  Q  GCAC	GACC+ CTTGG T CGTG GCAC V GGAC+ CCTG D GCAG CTTC Q GAAG	- 480 - 540 - - -6.0.0_				
a a	421 481 -541- 601	TTTCO K G AAGAA TTCTT K N GAGTO CTCAO E W TCCGA AGGCT S D GGGAA CCCTT	GGCA Q ACCAA CGGT Q GGGGA CCCT E ACGGG G CGCC G V CCTC	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCG. GGC' R CAGG S CAA' GTTC GTTC GTTC GAAG F CTTC GAAG F	AGA  E CCTTGGA  L TGGG  G ACC  G ATGGA  F ATGG  G ATGG  TTAC  C TCCC	ACC+ TGG P GAC'+ CTG' T GCA' CGT' Q CCT'+ GGA' S GGG' S	ACA- TGT- Q CTGGC- CGGC- CGGC- Y CGTA- GAT- GAT- V TAAA-	GGTCCTCCTCCTCCTCCTCCTCCTCCTCCTCCTCCTCCTC	GTAGE CATO	CAC( T GTG  T CAAL + GTT'  K CAAC T GTTC  L TGAC  L TGAC  TGAC CCC  TGAC  TG	CCT(CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	GCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCC. P CTA -+- SAT: Y SAC: -+- CCTG D GCA: CCTGT(	ATCO FINE S FICCO AGGO FINE STEEL T CAAC K K CAAC	CCGGC R CAGCC S S GCCC P GAGCC S GCCCGGA	D CGACO P CAGO R R CTACO R CTA	TGAC E ECATC I CGTC V GTGC W W CACC	GCTC  CGAC  L  CGGCGC  A  GCTC  CGAC  Q  GCAC	GACC+ CTTGG T CGTG GCAC V GGAC+ CCTG D GCAG CTTC Q GAAG	- 480 - 540 - - -6.0.0_				

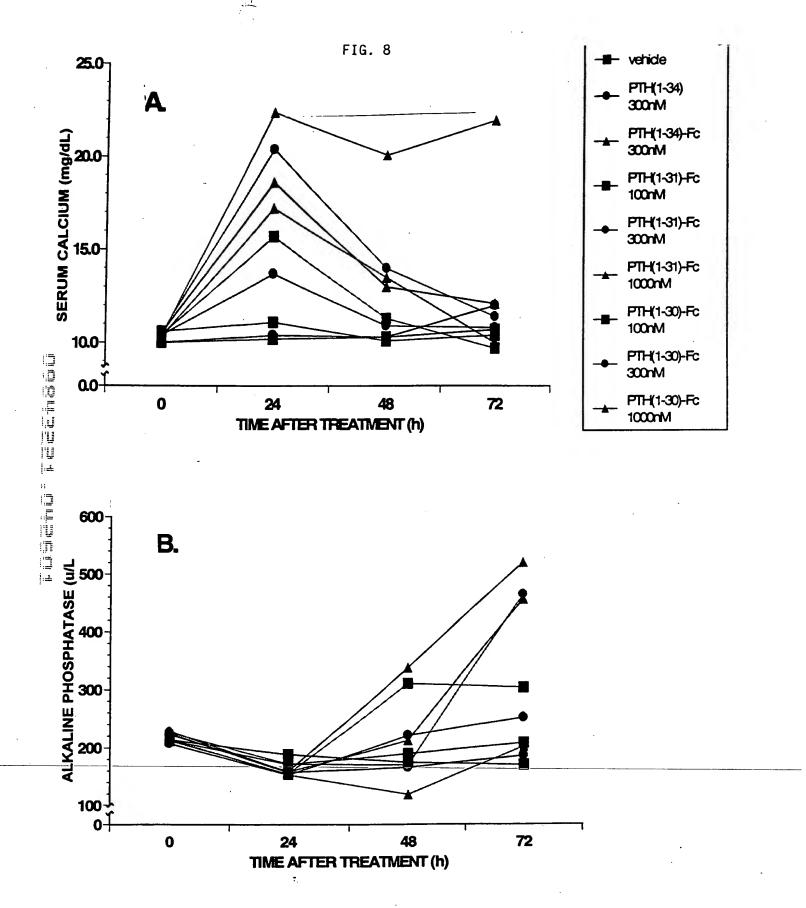


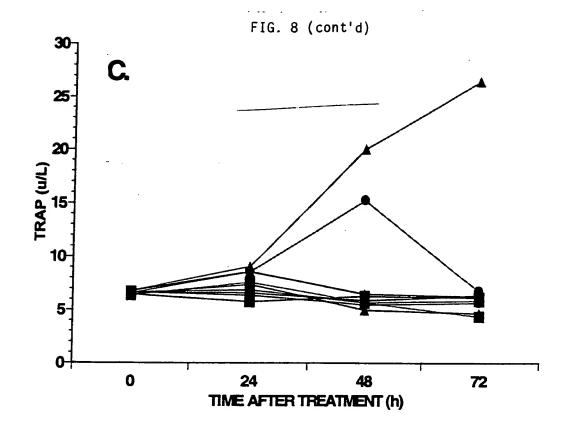


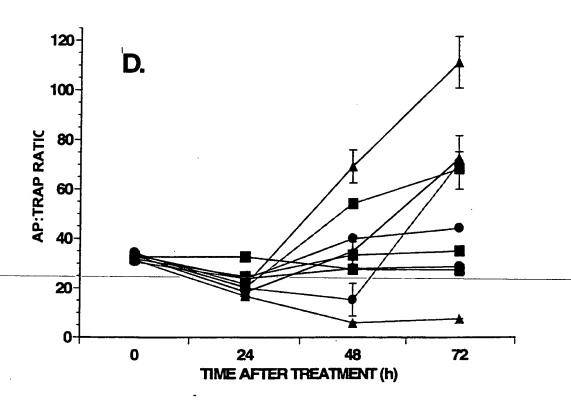


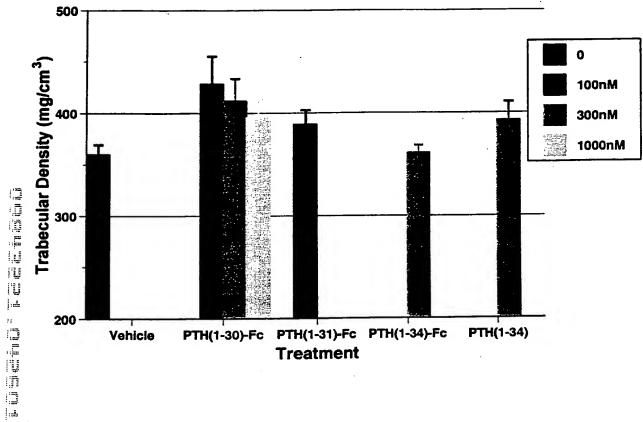












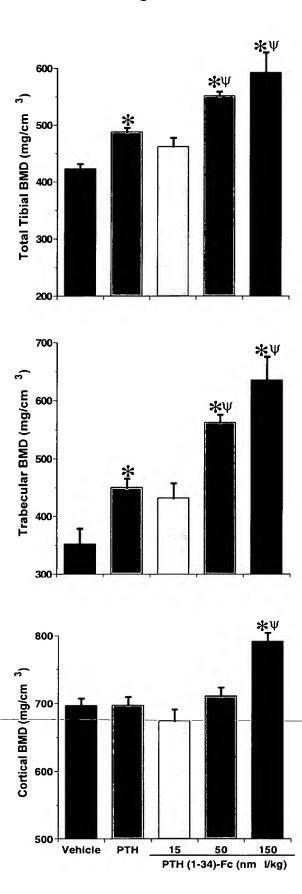
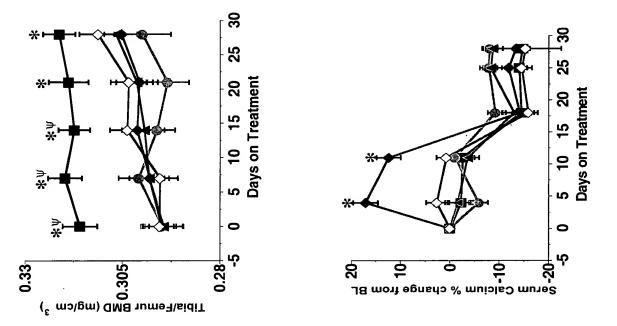
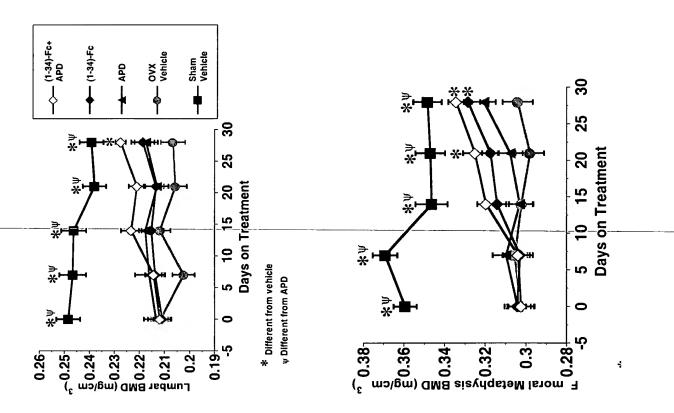


Fig. 11





## Effect of Single Dose SC Administration of $PTH_{(1-34)}Fc$ on Calcium

FIG. 12

